

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING**

### **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/562,324  
Source: IFWP  
Date Processed by STIC: 01/09/2006

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

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1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>**), EFS Submission User Manual - ePAVE)
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Revised 01/10/06



IFWO

## RAW SEQUENCE LISTING

DATE: 01/09/2006

PATENT APPLICATION: US/10/562,324

TIME: 10:43:50

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\01092006\J562324.raw

5 &lt;110&gt; APPLICANT: Bayer AG

9 &lt;120&gt; TITLE OF INVENTION: NEUES EIMERIA GEN UND PROTEIN SOWIE DEREN

## VERWENDUNG

13 &lt;130&gt; FILE REFERENCE: LeA 36695 DE

C--&gt; 17 &lt;140&gt; CURRENT APPLICATION NUMBER: US/10/562,324

C--&gt; 17 &lt;141&gt; CURRENT FILING DATE: 2005-12-23

17 &lt;160&gt; NUMBER OF SEQ ID NOS: 24

21 &lt;170&gt; SOFTWARE: PatentIn version 3.1

↓  
p/s do not use  
foreign language  
in US Application.

## ERRORED SEQUENCES

563 &lt;210&gt; SEQ ID NO: 24

565 &lt;211&gt; LENGTH: 18

567 &lt;212&gt; TYPE: DNA

569 &lt;213&gt; ORGANISM: Artificial Sequence

573 &lt;220&gt; FEATURE:

575 &lt;223&gt; OTHER INFORMATION: Primer BGH reverse

577 &lt;400&gt; SEQUENCE: 24

578 tagaaggcac agtcgagg

E--&gt; 584 Le a 36 695

E--&gt; 587 - 11 -

Does Not Comply  
Corrected Diskette Needed

(p3-1)

18

→ p/s delete

## VERIFICATION SUMMARY

DATE: 01/09/2006

PATENT APPLICATION: US/10/562,324

TIME: 10:43:51

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\01092006\J562324.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No  
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:45 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:41  
L:584 M:254 E: No. of Bases conflict, LENGTH:Input:695 Counted:21 SEQ:24  
L:584 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2  
L:584 M:112 C: (48) String data converted to lower case,  
L:587 M:254 E: No. of Bases conflict, this line has no nucleotides.  
L:587 M:252 E: No. of Seq. differs, <211> LENGTH:Input:18 Found:21 SEQ:24